

Does Not Comply
Corrected Diskette Needed
Errors were edited

1633

RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/356,575C

DATE: 11/16/2000
TIME: 15:08:51

Input Set : A:\pto.txt
Output Set: N:\CRF3\11162000\I356575C.raw

1 <110> APPLICANT: Fallaux, Frits Hoebe, ↓
2 Robert Bout, Abraham Valerio, Domenico van der Eb, Alex ↓
W--> 3 <120> TITLE OF INVENTION: PACKAGING SYSTEMS<130> 2578-3935US<140> ↓
W--> 4 <140> CURRENT APPLICATION NUMBER: US/09/356,575C
W--> 0 <130> FILE REFERENCE:
C--> 4 <141> CURRENT FILING DATE: 1999-07-19 ↓
5 <150> PRIOR APPLICATION NUMBER: 1995-06-14<160> EP 95201611.1<151> 1995-06-15<150> EP ↓
W--> 6 <151> PRIOR FILING DATE: 1995-06-26<160> 22 <170> PatentIn version 3.0<210> 1<211> ↓

ERRORED SEQUENCES

7 <212> TYPE: ~~DNA<213> Unknown<220><223>~~ Derived from Adenovirus
W--> 0 <160> NUMBER OF SEQ ID NOS:
W--> 9 <210> SEQ ID NO:
W--> 9 <211> LENGTH:
W--> 9 <213> ORGANISM:
E--> 9 <400> SEQUENCE: 1
E--> 10 cgtgtagtgt atttataccc g

21

Format errors.
Enter "hard
returns" where
arrows indicate.
These format
errors occur
throughout the
sequence listing.

Response and sequence counting errors also
due to the format errors cited above.

VERIFICATION SUMMARY

DATE: 11/16/2000

PATENT APPLICATION: US/09/356,575C

TIME: 15:08:52

Input Set : A:\pto.txt

Output Set : N:\CRF3\11162000\I356575C.raw

L:3 M:283 W: Missing Blank Line separator, <120> field identifier
L:4 M:282 W: Numeric Field Identifier Missing, <140> CURRENT APPLICATION NUMBER: is Added.
L:0 M:201 W: Mandatory field data missing, FILE REFERENCE
L:4 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:6 M:256 W: Invalid Numeric Header Field, Wrong Prior FILING DATE:YYYY-MM-DD
L:0 M:282 W: Numeric Field Identifier Missing, <160> is required.
L:9 M:282 W: Numeric Field Identifier Missing, <210> is required.
L:9 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:9 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:9 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:0 differs:1
L:10 M:252 E: No. of Seq. differs, <211>LENGTH:Input:0 Found:21 SEQ:0
L:15 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:15 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:15 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:21 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:21 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:21 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:27 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:27 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:27 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:33 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:33 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:33 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:39 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:39 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:39 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:45 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:45 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:45 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:51 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:51 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:51 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:57 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:57 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:57 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:63 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:63 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:63 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:69 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:69 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:69 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:75 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:75 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:75 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:81 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:81 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:81 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:87 M:282 W: Numeric Field Identifier Missing, <211> is required.

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Input Set : A:\pto.txt

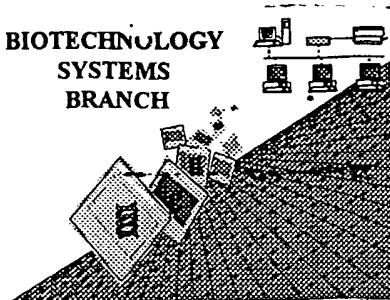
Output Set: N:\CRF3\11162000\I356575C.raw

L:87 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:87 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:93 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:93 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:93 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:99 M:282 W: Numeric Field Identifier Missing, <211> is required.
L:99 M:282 W: Numeric Field Identifier Missing, <212> is required.
L:99 M:282 W: Numeric Field Identifier Missing, <213> is required.
L:0 M:203 E: No. of Seq. differs, <160> Number Of Sequences:Input (0) Counted (22)

Wilson

RAW SEQUENCE LISTING ERROR REPORT

BIOTECHNOLOGY
SYSTEMS
BRANCH



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 09/356,575C
Source: 1633 RUSH
Date Processed by STIC: 10/31/2000

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

PATENTIN 3.0 e-mail help: patin30help@uspto.gov or phone 703-306-4119 (R. Wax)

→ TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.0 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW:

Checker Version 3.0

The Checker Version 3.0 application is a state-of-the-art Windows based software program employing a logical and intuitive user-interface to check whether a sequence listing is in compliance with format and content rules. Checker Version 3.0 works for sequence listings generated for the original version of 37 CFR §§1.821 - 1.825 effective October 1, 1990 (old rules) and the revised version (new rules) effective July 1, 1998 as well as World Intellectual Property Organization (WIPO) Standard ST.25.

Checker Version 3.0 replaces the previous DOS-based version of Checker, and is Y2K-compliant. Checker allows public users to check sequence listings in Computer Readable form (CRF) before submitting them to the United States Patent and Trademark Office (USPTO). Use of Checker prior to filing the sequence listing is expected to result in fewer errored sequence listings, thus saving time and money.

Checker Version 3.0 can be down loaded from the USPTO website at the following address:

<http://www.uspto.gov/web/offices/pac/checker>